

Appl. No. 09/833,328  
Amdt. dated March 14, 2005

PATENT

**APPENDIX A: REFORMATTED COPY OF ORIGINAL SEQUENCE LISTING**

## SEQUENCE LISTING

<110> Laemmle, Bernhard

Schwarz, Hans-Peter

Scheifflinger, Friedrich

Antoine, Gerhard

Kerschbaumer, Randolph

Tagliavacca, Luigina

Zimmermann, Klaus

Furlan, Miha

Turecek, Peter

Gerritsen, Helena E.

<120> Composition Exhibiting a von Willebrand Factor (vWF) Protease Activity  
Comprising a Polypeptide Chain with the Amino Acid Sequence AAGGILHLELLV

<130> 247.00CIP

<140> 09/833,328

<141> 2001-04-12

<150> 09/721,254

<151> 2000-11-22

<160> 15

<170> PatentIn version 3.1

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&lt;212&gt; PRT

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&lt;212&gt; PRT

&lt;213&gt; human

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Thr Asn Leu Asn Ile Gly Ala Glu Leu Leu Arg Asp Pro Ser Leu Gly  
20 25 30

Ala Gln Phe Arg Val His Leu Val Lys Met Val Ile Leu Thr Glu Pro  
35 40 45

Glu Gly Ala Pro Asn Ile Thr Ala Asn Leu Thr Ser Ser Leu Leu Ser  
50 55 60

Val Cys Gly Trp Ser Gln Thr Ile Asn Pro Glu Asp Asp Thr Asp Pro  
65 70 75 80

Gly His Ala Asp Leu Val Leu Tyr Ile Thr Arg Phe Asp Leu Glu Leu  
85 90 95

Pro Asp Gly Asn Arg Gln Val Arg Gly Val Thr Gln Leu Gly Gly Ala  
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Cys Ser Pro Thr Trp Ser Cys Leu Ile Thr Glu Asp Thr Gly Phe Asp  
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Leu Gly Val Thr Ile  
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cttoggggacc cgtccctggg ggctcagttt cgggtgcacc tgggtgaagat ggtcattctg      180
acagagcctg aggggtgccc aaatatcaca gccaacctca cctcgtccct gctgagcgtc      240
tgtgggtgga gccagaccat caaccctgag gacgacaagg atcctggcca tgotgacctg      300
gtcctotata tcaactaggtt tgacctggag ttgcctgatg gtaaccggca ggtgcggggc      360
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Asn Leu Asn Ile Gly Ala Glu Leu Leu Arg Asp Pro Ser Leu Gly Ala  
35 40 45

Gln Phe Arg Val His Leu Val Lys Met Val Ile Leu Thr Glu Pro Glu  
50 55 60

Gly Ala Pro Asn Ile Thr Ala Asp Leu Thr Ser Ser Leu Leu Ser Val  
65 70 75 80

Cys Gly Trp Ser Gln Thr Ile Asn Pro Glu Asp Asp Thr Asp Pro Gly  
85 90 95

His Ala Asp Leu Val Leu Tyr Ile Thr Arg Phe Asp Leu Glu Leu Pro  
100 105 110

Asp Gly Asn Arg Gln Val Arg Gly Val Thr Gln Leu Gly Gly Ala Cys  
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Gly Val Thr Ile  
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Leu Thr Asn Leu Asn Ile Gly Ala Glu Leu Leu Arg Asp Pro Ser Leu  
35 40 45  
Gly Ala Gln Phe Arg Val His Leu Val Lys Met Val Ile Leu Thr Glu  
50 55 60  
Pro Glu Gly Ala Pro Asn Ile Thr Ala Asn Leu Thr Ser Ser Leu Leu  
65 70 75 80  
Ser Val Cys Gly Trp Ser Gln Thr Ile Asn Pro Glu Asp Asp Thr Asp  
85 90 95  
Pro Gly His Ala Asp Leu Val Leu Tyr Ile Thr Arg Phe Asp Leu Glu  
100 105 110  
Leu Pro Asp Gly Asn Arg Gln Val Arg Gly Val Thr Gln Leu Gly Gly  
115 120 125  
Ala Cys Ser Pro Thr Trp Ser Cys Leu Ile Thr Glu Asp Thr Gly Phe  
130 135 140  
Asp Leu Gly Val Thr Ile  
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&lt;400&gt; 7

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<212> PRT

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<210> 15

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<212> PRT

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Ala Val Gly Pro Asp Val Phe Gln Ala His Gln Glu Asp Thr Glu Arg  
1 5 10 15

Tyr Val Leu Thr Asn Leu Asn Ile Gly Ala Glu Leu Leu Arg Asp Pro  
20 25 30

Ser Leu Gly Ala Gln Phe Arg Val His Leu Val Lys Met Val Ile Leu  
35 40 45

Thr Glu Pro Glu Gly Ala Pro Asn Ile Thr Ala Asn Leu Thr Ser Ser  
50 55 60

Leu Leu Ser Val Cys Gly Trp Ser Gln Thr Ile Asn Pro Glu Asp Asp  
65 70 75 80

Thr Asp Pro Gly His Ala Asp Leu Val Leu Tyr Ile Thr Arg Phe Asp  
85 90 95

Leu Glu Leu Pro Asp Gly Asn Arg Gln Val Arg Gly Val Thr Gln Leu  
100 105 110

Gly Gly Ala Cys Ser Pro Thr Trp Ser Cys Leu Ile Thr Glu Asp Thr  
115 120 125

Gly Phe Asp Leu Gly Val Thr Ile  
130 135